Serial Number: 09/ 645, 321

ENTERED

AU 1652
RECEIVED

- [ ] Changed a file from non-ASCII to ASCII
- [ ] Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- [ ] Edited a format error in the Current Application Data section, specifically: _____
- [ ] Edited the Current Application Data section with the actual current number. The number inputted by the applicant was [ ] the prior application data; or [ ] other _____.
- [ ] Added the mandatory heading and subheadings for "Current Application Data".
- [ ] Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- [ ] Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- [ ] Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- [ ] Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- [ ] Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- [ ] Inserted colons after headings/subheadings. Headings edited included: _____
- [ ] Deleted extra, invalid, headings used by an applicant, specifically: _____
- [ ] Deleted: [ ] non-ASCII "garbage" at the beginning/end of files; [ ] secretary initials/filename at end of file; [ ] page numbers throughout text; [ ] other invalid text, such as_____
- [ ] Inserted mandatory headings, specifically: _____
- [ ] Corrected an obvious error in the response, specifically: _____
- [ ] Edited identifiers where upper case is used but lower case is required, or vice versa.
- [ ] Corrected an error in the Number of Sequences field, specifically: _____
- [x] A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- [ ] Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- [ ] Other:
  _____
  _____
  _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.   3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/645,321

DATE: 09/14/2000
TIME: 16:50:58

Input Set : A:\Cpg.pto
Output Set: N:\CRF3\09142000\I645321.raw

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 3 <110> APPLICANT: Satoshi KOIZUMI
 4        Kazuhiko TABATA
 5        Tetsuo ENDO
 6        Akio OZAKI
 8 <120> TITLE OF INVENTION: Process for producing N-acetylneuraminic acid
10 <130> FILE REFERENCE: 11229
12 <140> CURRENT APPLICATION NUMBER: US/09/645,321
12 <141> CURRENT FILING DATE: 2000-08-25
12 <150> PRIOR APPLICATION NUMBER: H11-242670
13 <151> PRIOR FILING DATE: 1999-08-30
15 <160> NUMBER OF SEQ ID NOS: 8
17 <170> SOFTWARE: PatentIn Ver. 2.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 391
21 <212> TYPE: PRT
22 <213> ORGANISM: Synechocystis sp.(PCC6803)
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28 Leu His Gln Asp Val Leu Pro Phe Trp Glu Lys Tyr Ser Leu Asp Arg
29              20                  25                  30
31 Gln Gly Gly Gly Tyr Phe Thr Cys Leu Asp Arg Lys Gly Gln Val Phe
32              35                  40                  45
34 Asp Thr Asp Lys Phe Ile Trp Leu Gln Asn Arg Gln Val Trp Gln Phe
35          50                  55                  60
37 Ala Val Phe Tyr Asn Arg Leu Glu Pro Lys Pro Gln Trp Leu Glu Ile
38  65                  70                  75                  80
40 Ala Arg His Gly Ala Asp Phe Leu Ala Arg His Gly Arg Asp Gln Asp
41                  85                  90                  95
43 Gly Asn Trp Tyr Phe Ala Leu Asp Gln Glu Gly Lys Pro Leu Arg Gln
44              100                 105                 110
46 Pro Tyr Asn Val Phe Ser Asp Cys Phe Ala Ala Met Ala Phe Ser Gln
47              115                 120                 125
49 Tyr Ala Leu Ala Ser Gly Ala Gln Glu Ala Lys Ala Ile Ala Leu Gln
50          130                 135                 140
52 Ala Tyr Asn Asn Val Leu Arg Arg Gln His Asn Pro Lys Gly Gln Tyr
53 145                 150                 155                 160
55 Glu Lys Ser Tyr Pro Gly Thr Arg Pro Leu Lys Ser Leu Ala Val Pro
56              165                 170                 175
58 Met Ile Leu Ala Asn Leu Thr Leu Glu Met Glu Trp Leu Leu Pro Pro
59              180                 185                 190
61 Thr Thr Val Glu Glu Val Leu Ala Gln Thr Val Arg Glu Val Met Thr
62              195                 200                 205
64 Asp Phe Leu Asp Pro Glu Ile Gly Leu Met Arg Glu Ala Val Thr Pro
65          210                 215                 220
67 Thr Gly Glu Phe Val Asp Ser Phe Glu Gly Arg Leu Leu Asn Pro Gly
68 225                 230                 235                 240
```

Input Set : A:\Cpg.pto
Output Set: N:\CRF3\09142000\I645321.raw

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70 His Gly Ile Glu Ala Met Trp Phe Met Met Asp Ile Ala Gln Arg Ser
71                   215                 250                 255
73 Gly Asp Arg Gln Leu Gln Glu Gln Ala Ile Ala Val Val Leu Asn Thr
74                   260                 265                 270
76 Leu Glu Tyr Ala Trp Asp Glu Glu Phe Gly Gly Ile Phe Tyr Phe Leu
77           275                 280                 285
79 Asp Arg Gln Gly His Pro Pro Gln Gln Leu Glu Trp Asp Gln Lys Leu
80       290                 295                 300
82 Trp Trp Val His Leu Glu Thr Leu Val Ala Leu Ala Lys Gly His Gln
83 305                 310                 315                 320
85 Ala Thr Gly Gln Glu Lys Cys Trp Gln Trp Phe Glu Arg Val His Asp
86                   325                 330                 335
88 Tyr Ala Trp Ser His Phe Ala Asp Pro Glu Tyr Gly Glu Trp Phe Gly
89           340                 345                 350
91 Tyr Leu Asn Arg Arg Gly Glu Val Leu Leu Asn Leu Lys Gly Gly Lys
92           355                 360                 365
94 Trp Lys Gly Cys Phe His Val Pro Arg Ala Leu Trp Leu Cys Ala Glu
95       370                 375                 380
97 Thr Leu Gln Leu Pro Val Ser
98 385                 390
101 <210> SEQ ID NO: 2
102 <211> LENGTH: 1173
103 <212> TYPE: DNA
104 <213> ORGANISM: Synechocystis sp.(PCC6803)
106 <400> SEQUENCE: 2
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109  1              5                  10                  15
111 tta cac cag gac gta ttg ccc ttt tgg gaa aaa tat tcc ctc gat cgc   96
112 Leu His Gln Asp Val Leu Pro Phe Trp Glu Lys Tyr Ser Leu Asp Arg
113              20                  25                  30
115 cag ggg ggc ggt tac ttt acc tgc tta gac cgt aaa ggc cag gtt ttt  144
116 Gln Gly Gly Gly Tyr Phe Thr Cys Leu Asp Arg Lys Gly Gln Val Phe
117          35                  40                  45
119 gac aca gat aaa ttc att tgg tta caa aac cgt cag gta tgg cag ttt  192
120 Asp Thr Asp Lys Phe Ile Trp Leu Gln Asn Arg Gln Val Trp Gln Phe
121      50                  55                  60
123 gcc gtt ttc tac aac cgt ttg gaa cca aaa ccc caa tgg tta gaa att  240
124 Ala Val Phe Tyr Asn Arg Leu Glu Pro Lys Pro Gln Trp Leu Glu Ile
125  65                  70                  75                  80
127 gcc cgc cat ggt gct gat ttt tta gct cgc cac ggc cga gat caa gac  288
128 Ala Arg His Gly Ala Asp Phe Leu Ala Arg His Gly Arg Asp Gln Asp
129              85                  90                  95
131 ggt aat tgg tat ttt gct ttg gat cag gaa ggc aaa ccc ctg cgt caa  336
132 Gly Asn Trp Tyr Phe Ala Leu Asp Gln Glu Gly Lys Pro Leu Arg Gln
133          100                 105                 110
135 ccc tat aac gtt ttt tcc gat tgc ttc gcc gcc atg gcc ttt agt caa  384
136 Pro Tyr Asn Val Phe Ser Asp Cys Phe Ala Ala Met Ala Phe Ser Gln
137          115                 120                 125
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RAW SEQUENCE LISTING   DATE: 09/14/2000
PATENT APPLICATION   US/09/645,321   TIME: 16:50:58

Input Set : A:\Cpg.pto
Output Set: N:\CRF3\09142000\I645321.raw

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139 tat gcc tta gcc agt ggg gcg cag gaa gct aaa gcc att gcc ctg cag   432
140 Tyr Ala Leu Ala Ser Gly Ala Gln Glu Ala Lys Ala Ile Ala Leu Gln
141     130             135             140

143 gcc tac aat aac gtc cta cgc cgt cag cac aat ccc aaa ggt caa tac   480
144 Ala Tyr Asn Asn Val Leu Arg Arg Gln His Asn Pro Lys Gly Gln Tyr
145 115             150             155             160

147 gag aag tcc tat cca ggt act aga ccc ctc aaa tcc ctg gcg gtg ccg   528
148 Glu Lys Ser Tyr Pro Gly Thr Arg Pro Leu Lys Ser Leu Ala Val Pro
149             165             170             175

151 atg att tta gcc aac ctc acc ctg gag atg gaa tgg tta tta ccg cct   576
152 Met Ile Leu Ala Asn Leu Thr Leu Glu Met Glu Trp Leu Leu Pro Pro
153             180             185             190

155 act acc gtg gaa gag gtg ttg gcc caa acc gtc aga gaa gtg atg acg   624
156 Thr Thr Val Glu Glu Val Leu Ala Gln Thr Val Arg Glu Val Met Thr
157             195             200             205

159 gat ttc ctc gac cca gaa ata gga tta atg cgg gaa gcg gtg acc ccc   672
160 Asp Phe Leu Asp Pro Glu Ile Gly Leu Met Arg Glu Ala Val Thr Pro
161     210             215             220

163 aca gga gaa ttt gtt gat agt ttt gaa ggg cgg ttg ctc aac cca gga   720
164 Thr Gly Glu Phe Val Asp Ser Phe Glu Gly Arg Leu Leu Asn Pro Gly
165 225             230             235             240

167 cac ggc att gaa gcc atg tgg ttc atg atg gac att gcc caa cgc tcc   768
168 His Gly Ile Glu Ala Met Trp Phe Met Met Asp Ile Ala Gln Arg Ser
169             245             250             255

171 ggc gat cgc cag tta cag gag caa gcc att gca gtg gtg ttg aac acc   816
172 Gly Asp Arg Gln Leu Gln Glu Gln Ala Ile Ala Val Val Leu Asn Thr
173             260             265             270

175 ctg gaa tat gcc tgg gat gaa gaa ttt ggt ggc ata ttt tat ttc ctt   864
176 Leu Glu Tyr Ala Trp Asp Glu Glu Phe Gly Gly Ile Phe Tyr Phe Leu
177             275             280             285

179 gat cgc cag ggc cac cct ccc caa caa ctg gaa tgg gac caa aag ctc   912
180 Asp Arg Gln Gly His Pro Pro Gln Gln Leu Glu Trp Asp Gln Lys Leu
181     290             295             300

183 tgg tgg gta cat ttg gaa acc ctg gtt gcc cta gcc aag ggc cac caa   960
184 Trp Trp Val His Leu Glu Thr Leu Val Ala Leu Ala Lys Gly His Gln
185 305             310             315             320

187 gcc act ggc caa gaa aaa tgt tgg caa tgg ttt gag cgg gtc cat gat  1008
188 Ala Thr Gly Gln Glu Lys Cys Trp Gln Trp Phe Glu Arg Val His Asp
189             325             340             335

191 tac gcc tgg agt cat ttc gcc gat cct gag tat ggg gaa tgg ttt ggc  1056
192 Tyr Ala Trp Ser His Phe Ala Asp Pro Glu Tyr Gly Glu Trp Phe Gly
193             340             345             350

195 tac ctg aat cgc cgg gga gag gtg tta ctc aac cta aaa ggg ggg aaa  1104
196 Tyr Leu Asn Arg Arg Gly Glu Val Leu Leu Asn Leu Lys Gly Gly Lys
197             355             360             365

199 tgg aaa ggg tgc ttc cac gta ccc cga act ctg tgg ctc tgt gcg gaa  1152
200 Trp Lys Gly Cys Phe His Val Pro Arg Ala Leu Trp Leu Cys Ala Glu
201     370             375             380

203 act ctc caa ctt ccg gtt agt                                      1173
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RAW SEQUENCE LISTING   DATE: 09/14/2000
PATENT APPLICATION: US/09/645,321   TIME: 16:30:58

Input Set : A:\Cpg.pto
Output Set: N:\CRF3\09142000\I645321.raw

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205 385                    390
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209 <211> LENGTH: 24
210 <212> TYPE: DNA
211 <213> ORGANISM: Artificial Sequence
213 <220> FEATURE:
214 <223> OTHER INFORMATION: Synthetic DNA
216 <400> SEQUENCE: 3
217 gtgtaagctt tctgtatggg gtgt                            24
220 <210> SEQ ID NO: 4
221 <211> LENGTH: 26
222 <212> TYPE: DNA
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: Synthetic DNA
228 <400> SEQUENCE: 4
229 gcagggatcc caaccaggca gcggaa                          26
232 <210> SEQ ID NO: 5
233 <211> LENGTH: 32
234 <212> TYPE: DNA
235 <213> ORGANISM: Artificial Sequence
237 <220> FEATURE:
238 <223> OTHER INFORMATION: Synthetic DNA
240 <400> SEQUENCE: 5
241 tttatcgata ttaattaggg ggaatgaatg ag                   32
244 <210> SEQ ID NO: 6
245 <211> LENGTH: 33
246 <212> TYPE: DNA
247 <213> ORGANISM: Artificial Sequence
249 <220> FEATURE:
250 <223> OTHER INFORMATION: Synthetic DNA
252 <400> SEQUENCE: 6
253 tttggatcct cattattccc cctgattttt gaa                  33
256 <210> SEQ ID NO: 7
257 <211> LENGTH: 36
258 <212> TYPE: DNA
259 <213> ORGANISM: Artificial Sequence
261 <220> FEATURE:
262 <223> OTHER INFORMATION: Synthetic DNA
264 <400> SEQUENCE: 7
265 taaatcgata tttgtatgat tgcccatcgc cgtcag              36
268 <210> SEQ ID NO: 8
269 <211> LENGTH: 36
270 <212> TYPE: DNA
271 <213> ORGANISM: Artificial Sequence
273 <220> FEATURE:
274 <223> OTHER INFORMATION: Synthetic DNA
276 <400> SEQUENCE: 8
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9/14/00

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RAW SEQUENCE LISTING                    DATE: 09/14/2000
PATENT APPLICATION:   US/09/645,321     TIME: 16:30:58

Input Set  : A:\Cpg.pto
Output Set: N:\CRF3\09142000\I645321.raw
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277 aaaggatcct taactaaccg gaagttggag agtttc                    36
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VERIFICATION SUMMARY   DATE: 09/11/2000
PATENT APPLICATION: US/09/645,321   TIME: 16:50:59

Input Set : **A:\Cpg.pto**
Output Set: **N:\CRF3\09142000\I645321.raw**

L:12 M:270 C: Current Application Number differs. Replaced Current Application No
L:12 M:271 C: Current Filing Date differs. Replaced Current Filing Date

OIPE

RAW SEQUENCE LISTING   DATE: 09/06/2000
PATENT APPLICATION: US/09/645,321   TIME: 11:55:35

Input Set : A:\5.1183 sequence.txt
Output Set: N:\CRF3\09062000\I645321.raw

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    3 <110> APPLICANT: Satoshi KOIZUMI
    4         Kazuhiko TABATA
    5         Tetsuo ENDO
    6         Akio OZAKI
    8 <120> TITLE OF INVENTION: Process for producing N-acetylneuraminic acid
   10 <130> FILE REFERENCE: 11229
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/645,321
C--> 12 <141> CURRENT FILING DATE: 2000-08-25
   12 <150> PRIOR APPLICATION NUMBER: H11-242670
   13 <151> PRIOR FILING DATE: 1999-08-30
   15 <160> NUMBER OF SEQ ID NOS: 8
   17 <170> SOFTWARE: PatentIn Ver. 2.0
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ERRORED SEQUENCES

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     101 <210> SEQ ID NO: 2
     102 <211> LENGTH: 1173
     103 <212> TYPE: DNA
     101 <213> ORGANISM: Synechocystis sp.(PCC6803)
     106 <400> SEQUENCE: 2
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E--> 108 48
     109 Met Ile Ala His Arg Arg Gln Glu Leu Ala Gln Gln Tyr Tyr Gln Ala
     110  1               5                  10                  15
E--> 112 tta cac cag gac gta ttg ccc ttt tgg gaa aaa tat tcc ctc gat cgc
E--> 113 96
     114 Leu His Gln Asp Val Leu Pro Phe Trp Glu Lys Tyr Ser Leu Asp Arg
     115              20                  25                  30
E--> 117 cag ggg ggc ggt tac ttt acc tgc tta gac cgt aaa ggc cag gtt ttt
E--> 118 144
     119 Gln Gly Gly Gly Tyr Phe Thr Cys Leu Asp Arg Lys Gly Gln Val Phe
     120          35                  10                  45
E--> 122 gac aca gat aaa ttc att tgg tta caa aac cgt cag gta tgg cag ttt
E--> 123 192
     121 Asp Thr Asp Lys Phe Ile Trp Leu Gln Asn Arg Gln Val Trp Gln Phe
     125      50                  55                  60
E--> 127 gcc gtt ttc tac aac cgt ttg gaa cca aaa ccc caa tgg tta gaa att
E--> 128 240
     129 Ala Val Phe Tyr Asn Arg Leu Glu Pro Lys Pro Gln Trp Leu Glu Ile
     130  65                  70                  75                  80
E--> 132 gcc cgc cat ggt gct gat ttt tta gct cgc cac ggc cga gat caa gac
E--> 133 288
     134 Ala Arg His Gly Ala Asp Phe Leu Ala Arg His Gly Arg Asp Gln Asp
     135              85                  90                  95
E--> 137 ggt aat tgg tat ttt gct ttg gat cag gaa ggc aaa ccc ctg cgt caa
E--> 138 336
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[Handwritten annotations:]

Does Not Comply
Corrected Diskette Needed

Nucleic acid numbers "wrapped" down to the next line Possible "Hard page break" error

RAW SEQUENCE LISTING   DATE: 09/06/2000
PATENT APPLICATION: US/09/645,321   TIME: 11:55:35

Input Set : A:\5.1183 sequence.txt
Output Set: N:\CRF3\09062000\I645321.raw

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      139 Gly Asn Trp Tyr Phe Ala Leu Asp Gln Glu Gly Lys Pro Leu Arg Gln
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E--> 142 ccc tat aac gtt ttt tcc gat tgc ttc gcc gcc atg gcc ttt agt caa
E--> 143 384
      144 Pro Tyr Asn Val Phe Ser Asp Cys Phe Ala Ala Met Ala Phe Ser Gln
      115         115             120             125
E--> 147 tat gcc tta gcc agt ggg gcg cag gaa gct aaa gcc att gcc ctg cag
E--> 148 432
      149 Tyr Ala Leu Ala Ser Gly Ala Gln Glu Ala Lys Ala Ile Ala Leu Gln
      150     130             135             140
E--> 152 gcc tac aat aac gtc cta cgc cgt cag cac aat ccc aaa ggt caa tac
E--> 153 480
      154 Ala Tyr Asn Asn Val Leu Arg Arg Gln His Asn Pro Lys Gly Gln Tyr
      155 145             150             155             160
E--> 157 gag aag tcc tat cca ggt act aga ccc ctc aaa tcc ctg gcg gtg ccg
E--> 158 528
      159 Glu Lys Ser Tyr Pro Gly Thr Arg Pro Leu Lys Ser Leu Ala Val Pro
      160             165             170             175
E--> 162 atg att tta gcc aac ctc acc ctg gag atg gaa tgg tta tta ccg cct
E--> 163 576
      164 Met Ile Leu Ala Asn Leu Thr Leu Glu Met Glu Trp Leu Leu Pro Pro
      165             180             185             190
E--> 167 act acc gtg gaa gag gtg ttg gcc caa acc gtc aga gaa gtg atg acg
E--> 168 624
      169 Thr Thr Val Glu Glu Val Leu Ala Gln Thr Val Arg Glu Val Met Thr
      170         195             200             205
E--> 172 gat ttc ctc gac cca gaa ata gga tta atg cgg gaa gcg gtg acc ccc
E--> 173 672
      174 Asp Phe Leu Asp Pro Glu Ile Gly Leu Met Arg Glu Ala Val Thr Pro
      175     210             215             220
E--> 177 aca gga gaa ttt gtt gat agt ttt gaa ggg cgg ttg ctc aac cca gga
E--> 178 720
      179 Thr Gly Glu Phe Val Asp Ser Phe Glu Gly Arg Leu Leu Asn Pro Gly
      180 225             230             235             240
E--> 182 cac ggc att gaa gcc atg tgg ttc atg atg gac att gcc caa cgc tcc
E--> 183 768
      184 His Gly Ile Glu Ala Met Trp Phe Met Met Asp Ile Ala Gln Arg Ser
      185             245             250             255
E--> 187 ggc gat cgc cag tta cag gag caa gcc att gca gtg gtg ttg aac acc
E--> 188 816
      189 Gly Asp Arg Gln Leu Gln Glu Gln Ala Ile Ala Val Val Leu Asn Thr
      190         260             265             270
E--> 192 ctg gaa tat gcc tgg gat gaa gaa ttt ggt ggc ata ttt tat ttc ctt
E--> 193 864
      194 Leu Glu Tyr Ala Trp Asp Glu Glu Phe Gly Gly Ile Phe Tyr Phe Leu
      195         275             280             285
E--> 197 gat cgc cag ggc cac cct ccc caa caa ctg gaa tgg gac caa aag ctc
E--> 198 912
      199 Asp Arg Gln Gly His Pro Pro Gln Gln Leu Glu Trp Asp Gln Lys Leu
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RAW SEQUENCE LISTING   DATE: 09/06/2000
PATENT APPLICATION   US/09/645,321   TIME: 11:35:35

Input Set : A:\5.1183 sequence.txt
Output Set: N:\CRF3\09062000\I645321.raw

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        200      290              295            300
E--> 202 tgg tgg gta cat ttg gaa acc ctg gtt gcc cta gcc aag ggc cac caa
E--> 203 960
     204 Trp Trp Val His Leu Glu Thr Leu Val Ala Leu Ala Lys Gly His Gln
     205 305        310                 315                320
E--> 207 gcc act ggc caa gaa aaa tgt tgg caa tgg ttt gag cgg gtc cat gat
E--> 208 1008
     209 Ala Thr Gly Gln Glu Lys Cys Trp Gln Trp Phe Glu Arg Val His Asp
     210        325              330                335
E--> 212 tac gcc tgg agt cat ttc gcc gat cct gag tat ggg gaa tgg ttt ggc
E--> 213 1056
     211 Tyr Ala Trp Ser His Phe Ala Asp Pro Glu Tyr Gly Glu Trp Phe Gly
     215        340              345                350
E--> 217 tac ctg aat cgc cgg gga gag gtg tta ctc aac cta aaa ggg ggg aaa
E--> 218 1104
     219 Tyr Leu Asn Arg Arg Gly Glu Val Leu Leu Asn Leu Lys Gly Gly Lys
     220        355              360              365
E--> 222 tgg aaa ggg tgc ttc cac gtg ccc cga gct ctg tgg ctc tgt gcg gaa
E--> 223 1152
     224 Trp Lys Gly Cys Phe His Val Pro Arg Ala Leu Trp Leu Cys Ala Glu
     225        370              375              380
E--> 227 act ctc caa ctt ccg gtt agt
E--> 228 1173
     229 Thr Leu Gln Leu Pro Val Ser
     230 385              390
     233 <210> SEQ ID NO: 3
     234 <211> LENGTH: 24
     235 <212> TYPE: DNA
     236 <213> ORGANISM: Artificial Sequence
     238 <220> FEATURE:
     239 <223> OTHER INFORMATION: Synthetic DNA
     211 <400> SEQUENCE: 3
E--> 242 gtgtaagctt tctgtatggg gtgt
     243        24
     246 <210> SEQ ID NO: 1
     247 <211> LENGTH: 26
     248 <212> TYPE: DNA
     249 <213> ORGANISM: Artificial Sequence
     251 <220> FEATURE:
     252 <223> OTHER INFORMATION: Synthetic DNA
     254 <400> SEQUENCE: 1
E--> 255 gcaggatcc caaccaggca gcggaa
     256        26
     259 <210> SEQ ID NO: 5
     260 <211> LENGTH: 32
     261 <212> TYPE: DNA
     262 <213> ORGANISM: Artificial Sequence
     264 <220> FEATURE:
     265 <223> OTHER INFORMATION: Synthetic DNA
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refer to p.1

Same, p.1

Same, p.1

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/645,321

DATE: 09/06/2000
TIME: 11:55:35

Input Set : A:\5.1183 sequence.txt
Output Set: N:\CRF3\09062000\I645321.raw

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E--> 268 tttatcgata ttaattaggg ggaatgaatg ag
         269       32
         272  <210> SEQ ID NO: 6
         273  <211> LENGTH: 33
         274  <212> TYPE: DNA
         275  <213> ORGANISM: Artificial Sequence
         277  <220> FEATURE:
         278  <223> OTHER INFORMATION: Synthetic DNA
         280  <400> SEQUENCE: 6
E--> 281 tttggatcct cattattccc cctgattttt gaa
         282       33
         285  <210> SEQ ID NO: 7
         286  <211> LENGTH: 36
         287  <212> TYPE: DNA
         288  <213> ORGANISM: Artificial Sequence
         290  <220> FEATURE:
         291  <223> OTHER INFORMATION: Synthetic DNA
         293  <400> SEQUENCE: 7
E--> 294 taaatcgata tttgtatgat tgcccatcgc cgtcag
E--> 295 36
         298  <210> SEQ ID NO: 8
         299  <211> LENGTH: 36
         300  <212> TYPE: DNA
         301  <213> ORGANISM: Artificial Sequence
         303  <220> FEATURE:
         304  <223> OTHER INFORMATION: Synthetic DNA
         306  <400> SEQUENCE: 8
E--> 307 aaaggatcct taactaaccg gaagttggag agtttc
E--> 308 36
```

Same refer to p.1

9/6/00

file://C:\CRF3\Outhold\VsrI645321.htm